## GenCore version 5.1.7 Copyright (c) 1993 - 2006 Biocceleration Ltd.

OM nucleic - nucleic search, using sw model

Run on: April 14, 2006, 16:49:04; Search time 363 Seconds

(without alignments)

9470.535 Million cell updates/sec

Title: US-10-086-623-5

Perfect score: 1934

Sequence: 1 ttgtaccgaagagatgagac.....atcgacgtaactggaaaccg 1934

Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapext 1.0

Searched: 1303057 seqs, 888780828 residues

Total number of hits satisfying chosen parameters: 2606114

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents\_NA:\*

1: /cgn2\_6/ptodata/1/ina/1\_COMB.seq:\*

2: /cgn2\_6/ptodata/1/ina/5\_COMB.seq:\*

3: /cgn2\_6/ptodata/1/ina/6A\_COMB.seq:\*

4: /cgn2\_6/ptodata/1/ina/6B\_COMB.seq:\*

5: /cgn2\_6/ptodata/1/ina/H\_COMB.seq:\*

6: /cgn2\_6/ptodata/1/ina/PCTUS\_COMB.seq:\*

7: /cgn2 6/ptodata/1/ina/PP COMB.seq:\*

8: /cgn2\_6/ptodata/1/ina/RE\_COMB.seq:\*

9: /cgn2 6/ptodata/1/ina/backfiles1.seq:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

Result		% Query Match	Length	DB			
No.	Score				ID	Description	
1	1934	100.0	1934	3	US-09-438-046-5	Sequence 5, Appli	
2	1934	100.0	2253	3	US-09-438-046-7	Sequence 7, Appli	
3	1491.8	77.1	1882	3	US-09-457-066-36	Sequence 36, Appl	
4	1491.8	77.1	1882	3	US-09-540-224-1	Sequence 1, Appli	
5	1491.8	77.1	1882	3	US-09-564-595D-1	Sequence 1, Appli	
6	1491.8	77.1	1882	3	US-09-706-968-36	Sequence 36, Appl	
7	1491.8	77.1	1882	3	US-09-808-972-1	Sequence 1, Appli	
8	1491.8	77.1	1882	3	US-10-139-583-36	Sequence 36, Appl	

```
9
      1491.8
                77.1
                       1882 3
                                US-10-039-847A-1
                                                            Sequence 1, Appli
   10
      1491.8
                77.1
                       1882
                             3
                                US-10-226-559-1
                                                           Sequence 1, Appli
   11 1491.8
                77.1
                       1882 3
                                US-09-541-752-36
                                                           Sequence 36, Appl
   12
       1491.8
                77.1
                       1882 3
                                                           Sequence 1, Appli
                                US-09-876-813A-1
   13
        795.4
                41.1
                       1472 3
                                US-09-540-224-3
                                                            Sequence 3, Appli
   14
        795.4
                41.1
                       1472 3
                                US-09-564-595D-52
                                                           Sequence 52, Appl
   15
        795.4
                41.1
                       1472 3
                                US-09-808-972-3
                                                           Sequence 3, Appli
                       1472 3
   16
        795.4
                41.1
                                US-10-039-847A-3
                                                           Sequence 3, Appli
   17
        795.4
                41.1
                       1472 3
                                US-10-226-559-3
                                                           Sequence 3, Appli
   18
        795.4
                41.1
                       1472 3
                                US-09-876-813A-52
                                                            Sequence 52, Appl
   19
        697.6
                36.1
                       1110 3
                                US-09-564-595D-6
                                                            Sequence 6, Appli
   20
        697.6
                36.1
                       1110 3
                                US-09-876-813A-6
                                                            Sequence 6, Appli
   21
          662
                34.2
                        690 3
                                US-09-438-046-3
                                                            Sequence 3, Appli
        355.2
   22
                18.4
                        360 3
                                                            Sequence 1, Appli
                                US-09-438-046-1
  23
                        256
        213.8
                11.1
                             3
                                US-09-222-575-13
                                                            Sequence 13, Appl
        213.8
                11.1
                        256
                             3
                                US-09-389-681-13
С
  24
                                                            Sequence 13, Appl
c 25
        213.8
                11.1
                        256 3
                                US-09-620-405B-13
                                                            Sequence 13, Appl
c 26
        213.8
                11.1
                        256 3
                                US-09-339-338-13
                                                            Sequence 13, Appl
c 27
        213.8
                11.1
                        256 3
                                US-09-433-826B-13
                                                            Sequence 13, Appl
c 28
        213.8
                11.1
                        256 3
                                US-09-604-287A-13
                                                            Sequence 13, Appl
                        256 3
С
  29
        213.8
                11.1
                                US-09-285-480-13
                                                            Sequence 13, Appl
                        256 3
С
  30
        213.8
                11.1
                                US-09-834-759-13
                                                            Sequence 13, Appl
                                                            Sequence 13, Appl
С
  31
        213.8
                11.1
                        256 3
                                US-09-590-751A-13
С
  32
        213.8
                11.1
                        256 3
                                US-09-551-621-13
                                                            Sequence 13, Appl
С
  33
        213.8
                11.1
                        256 3
                                US-09-551-621A-13
                                                            Sequence 13, Appl
        213.8
                11.1
                        256 3
                                                            Sequence 13, Appl
  34
                                US-10-076-622-13
   35
        181.8
                 9.4
                       1035 3
                                US-09-457-066-6
                                                            Sequence 6, Appli
                       1035 3
   36
        181.8
                 9.4
                                US-09-706-968-6
                                                            Sequence 6, Appli
   37
        181.8
                 9.4
                       1035
                             3
                                US-10-139-583-6
                                                            Sequence 6, Appli
   38
        181.8
                 9.4
                       1035 3
                                US-09-541-752-6
                                                            Sequence 6, Appli
   39
        123.2
                 6.4
                       3571 3
                                US-09-457-066-42
                                                            Sequence 42, Appl
   40
        123.2
                 6.4
                       3571 3
                                US-09-564-595D-34
                                                            Sequence 34, Appl
   41
                       3571 3
                                US-09-706-968-42
                                                            Sequence 42, Appl
        123.2
                 6.4
   42
        123.2
                       3571 3
                                US-09-823-033-3
                                                            Sequence 3, Appli
                 6.4
                       3571 3
   43
        123.2
                 6.4
                                US-10-139-583-42
                                                            Sequence 42, Appl
                       3571
   44
        123.2
                 6.4
                             3
                                US-09-541-752-42
                                                            Sequence 42, Appl
                       3571 3
   45
        123.2
                 6.4
                                US-09-695-121-3
                                                            Sequence 3, Appli
```

#### ALIGNMENTS

```
; Sequence 5, Application US/09438046
; Patent No. 6706687
; GENERAL INFORMATION:
; APPLICANT: ERIKSSON, Ulf
; APPLICANT: AASE, Karin
; APPLICANT: LEE, Xuri
; APPLICANT: PONTN, Annica
; APPLICANT: UUTELA, Marko
; APPLICANT: ALITALO, Kari
; APPLICANT: OESTMAN, Arne
; APPLICANT: HELDIN, Carl-Henrik
; TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR D, DNA CODING
; TITLE OF INVENTION: THEREFOR, AND USES THEREOF
```

RESULT 1

US-09-438-046-5

### GenCore version 5.1.7 Copyright (c) 1993 - 2006 Biocceleration Ltd.

OM nucleic - protein search, using frame\_plus\_n2p model

Ruń on: April 14, 2006, 14:31:10; Search time 12.9 Seconds

(without alignments)

2478.989 Million cell updates/sec

Title: US-10-086-623-5

Perfect score: 3446

Sequence: 1 ttgtaccgaagagatgagac.....atcgacgtaactggaaaccg 1934

Scoring table: BLOSUM62

Xgapop 10.0 , Xgapext 0.5 Ygapop 10.0 , Ygapext 0.5 Fgapop 6.0 , Fgapext 7.0 Delop 6.0 , Delext 7.0

Searched: 572060 segs, 82675679 residues

Total number of hits satisfying chosen parameters: 1144120

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

### Command line parameters:

- -MODEL=frame+ n2p.model -DEV=xlp
- -Q=/abss/ABSSWEB\_spool/US10086623/runat\_14042006\_123917\_255/app\_query.fasta\_1
- -DB=Issued Patents AA -QFMT=fastan -SUFFIX=rai -MINMATCH=0.1 -LOOPCL=0
- -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
- -LIST=45 -DOCALIGN=200 -THR\_SCORE=pct -THR\_MAX=100 -THR\_MIN=0 -ALIGN=15
- -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
- -HOST=abss02p -USER=US10086623 @CGN 1\_1\_71\_@runat\_14042006\_123917\_255 -NCPU=6
- -ICPU=3 -NO MMAP -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV TIMEOUT=120
- -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
- -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7

Database : Issued Patents AA:\*

- 1: /cgn2\_6/ptodata/1/iaa/5\_COMB.pep:\*
- 2: /cgn2\_6/ptodata/1/iaa/6\_COMB.pep:\*
- 3: /cgn2 6/ptodata/1/iaa/H COMB.pep:\*
- 4: /cgn2\_6/ptodata/1/iaa/PCTUS\_COMB.pep:\*
- 5: /cgn2\_6/ptodata/1/iaa/RE\_COMB.pep:\*
- 6: /cgn2\_6/ptodata/1/iaa/backfiles1.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Dogult		ð Ou orst				
Result	Caoro	Query	Tonath I	מכ	TD	Doggrintion
No.	Score	Match	Length I	פע	ID	Description
1	1740	50.6	322	2	US-09-438-046-6	Sequence 6, Appli
1 2	1742	50.6	370	2	US-09-457-066-37	Sequence 37, Appl
	1742					Sequence 2, Appli
3	1742	50.6	370	2	US-09-540-224-2	
4	1742	50.6	370	2	US-09-564-595D-2	Sequence 2, Appli
5	1742	50.6	370	2	US-09-706-968-37	Sequence 37, Appl
6	1742	50.6	370	2	US-09-808-972-2	Sequence 2, Appli
7	1742	50.6	370	2	US-09-823-033-5	Sequence 5, Appli
8	1742	50.6	370	2	US-09-438-046-8	Sequence 8, Appli
9	1742	50.6	370	2	US-10-139-583-37	Sequence 37, Appl
. 10	1742	50.6	370	2	US-10-039-847A-2	Sequence 2, Appli
11	1742	50.6	370	2	US-10-226-559-2	Sequence 2, Appli
12	1742	50.6	370	2	US-09-541-752-37	Sequence 37, Appl
13	1742	50.6	370	2	US-09-695-121-5	Sequence 5, Appli
14	1742	50.6	370	2	US-09-876-813A-2	Sequence 2, Appli
15	1561	45.3	370	2	US-09-540-224-4	Sequence 4, Appli
16	1561	45.3	370	2	US-09-564-595D-53	Sequence 53, Appl
17	1561	45.3	370	2	US-09-808-972-4	Sequence 4, Appli
18	1561	45.3	370	2	US-10-039-847A-4	Sequence 4, Appli
19	1561	45.3	370	2	US-10-226-559-4	Sequence 4, Appli
20	1561	45.3	370	2	US-09-876-813A-53	Sequence 53, Appl
21	1403	40.7	317	2	US-09-564-595D-56	Sequence 56, Appl
22	1357.5	39.4	316	2	US-09-564-595D-55	Sequence 55, Appl
23	1095	31.8	303	2	US-09-564-595D-57	Sequence 57, Appl
24	1061.5	30.8	302	2	US-09-564-595D-54	Sequence 54, Appl
25	1037	30.1	200	2	US-09-438-046-4	Sequence 4, Appli
26	750.5	21.8	345	2	US-09-457-066-43	Sequence 43, Appl
27	750.5	21.8	345	2	US-09-564-595D-35	Sequence 35, Appl
28	750.5	21.8	345	2	US-09-706-968-43	Sequence 43, Appl
29	750.5	21.8	345	2	US-09-823-033-4	Sequence 4, Appli
30	750.5	21.8	345	2	US-10-139-583-43	Sequence 43, Appl
31	750.5	21.8	345	2	US-09-541-752-43	Sequence 43, Appl
32	750.5	21.8	345	2	US-09-695-121-4	Sequence 4, Appli
33	750.5	21.8	345	2	US-09-876-813A-35	Sequence 35, Appl
34	741	21.5	374	2	US-09-468-647A-118	Sequence 118, App
35	739.5	21.5	323	2	US-09-468-647A-1	Sequence 1, Appli
36	739.5	21.5	345	2	US-09-040-220D-2	Sequence 2, Appli
37	739.5	21.5	345	2	US-09-457-066-2	Sequence 2, Appli
38	739.5	21.5	345	2	US-09-265-686-2	Sequence 2, Appli
39	739.5	21.5	345	2	US-09-540-224-5	Sequence 5, Appli
40	739.5	21.5	345	2	US-09-564-595D-33	Sequence 33, Appl
41	739.5	21.5	345	2	US-09-706-968-2	Sequence 2, Appli
42	739.5	21.5	345	2	US-09-723-749-2	Sequence 2, Appli
43	739.5	21.5	345	2	US-09-823-033-2	Sequence 2, Appli
44	739.5	21.5		2	US-09-468-647A-2	Sequence 2, Appli
45	739.5	21.5		2	US-09-468-647A-101	Sequence 101, App
43	139.3		243	~	00 00 100 01/11 101	,pp

# ALIGNMENTS

## RESULT 1 US-09-438-046-6

<sup>;</sup> Sequence 6, Application US/09438046 ; Patent No. 6706687

```
; GENERAL INFORMATION:
  APPLICANT: ERIKSSON, Ulf
  APPLICANT: AASE, Karin
  APPLICANT: LEE, Xuri
  APPLICANT: PONTN, Annica
  APPLICANT: UUTELA, Marko
  APPLICANT: ALITALO, Kari
  APPLICANT: OESTMAN, Arne
  APPLICANT: HELDIN, Carl-Henrik
  TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR D,
                                                    DNA CODING
  TITLE OF INVENTION: THEREFOR, AND USES THEREOF
  FILE REFERENCE: Ulf Eriksson et al
  CURRENT APPLICATION NUMBER: US/09/438,046
  CURRENT FILING DATE: 1999-11-10
  EARLIER APPLICATION NUMBER: 60/107,852
  EARLIER FILING DATE: 1998-11-10
  EARLIER APPLICATION NUMBER: 60/113,997
  EARLIER FILING DATE: 1999-12-28
  EARLIER APPLICATION NUMBER: 60/150,604
  EARLIER FILING DATE: 1999-08-26
  EARLIER APPLICATION NUMBER: 60/157,108
  EARLIER FILING DATE: 1999-10-04
  EARLIER APPLICATION NUMBER: 60/157,756
  EARLIER FILING DATE: 1999-10-05
  NUMBER OF SEQ ID NOS: 31
  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 6
   LENGTH: 322
   TYPE: PRT
   ORGANISM: Homo sapiens
US-09-438-046-6
Alignment Scores:
Pred. No.:
                                   Length:
                                                322
                     4.43e-191
                                                322
                                   Matches:
Score:
                     1742.00
Percent Similarity:
                     100.0%
                                   Conservative:
                     100.0%
                                   Mismatches:
                                                0
Best Local Similarity:
                     50.6%
                                   Indels:
                                                0
Query Match:
                                                0
                                   Gaps:
DB:
US-10-086-623-5 (1-1934) x US-09-438-046-6 (1-322)
          1 TTGTACCGAAGAGATGAGACCATCCAGGTGAAAGGAAACGGCTACGTGCAGAGTCCTAGA 60
Qу
            1 LeuTyrArgArgAspGluThrIleGlnValLysGlyAsnGlyTyrValGlnSerProArg .20
Db
          61 TTCCCGAACAGCTACCCCAGGAACCTGCTCCTGACATGGCGGCTTCACTCTCAGGAGAAT 120
Qу
            21 PheProAsnSerTyrProArgAsnLeuLeuLeuThrTrpArgLeuHisSerGlnGluAsn 40
Db
         121 ACACGGATACAGCTAGTGTTTGACAATCAGTTTGGATTAGAGGAAGCAGAAAATGATATC 180
Qy
            41 ThrArgIleGlnLeuValPheAspAsnGlnPheGlyLeuGluGluAlaGluAsnAspIle 60
Db
         181 TGTAGGTATGATTTTGTGGAAGTTGAAGATATATCCGAAACCAGTACCATTATTAGAGGA 240
Qу
             61 CysArgTyrAspPheValGluValGluAspIleSerGluThrSerThrIleIleArgGly 80
Db
```

```
Qy
          81 ArgTrpCysGlyHisLysGluValProProArgIleLysSerArgThrAsnGlnIleLys 100
Db
       301 ATCACATTCAAGTCCGATGACTACTTTGTGGCTAAACCTGGATTCAAGATTTATTATTCT 360
Qy
          101 IleThrPheLysSerAspAspTyrPheValAlaLysProGlyPheLysIleTyrTyrSer 120
Db
       361 TTGCTGGAAGATTTCCAACCCGCAGCAGCTTCAGAGACCAACTGGGAATCTGTCACAAGC 420
Qу
          121 LeuLeuGluAspPheGlnProAlaAlaAlaSerGluThrAsnTrpGluSerValThrSer 140
Db
       421 TCTATTTCAGGGGTATCCTATAACTCTCCATCAGTAACGGATCCCACTCTGATTGCGGAT 480
Qy
          141 SerIleSerGlyValSerTyrAsnSerProSerValThrAspProThrLeuIleAlaAsp 160
Db
       481 GCTCTGGACAAAAAATTGCAGAATTTGATACAGTGGAAGATCTGCTCAAGTACTTCAAT 540
Qу
         161 AlaLeuAspLysLysIleAlaGluPheAspThrValGluAspLeuLeuLysTyrPheAsn 180
Db
       541 CCAGAGTCATGGCAAGAAGATCTTGAGAATATGTATCTGGACACCCCTCGGTATCGAGGC 600
Qу
          181 ProGluSerTrpGlnGluAspLeuGluAsnMetTyrLeuAspThrProArgTyrArgGly 200
Db
       601 AGGTCATACCATGACCGGAAGTCAAAAGTTGACCTGGATAGGCTCAATGATGATGCCAAG 660
Qу
          201 ArgSerTyrHisAspArgLysSerLysValAspLeuAspArgLeuAsnAspAspAlaLys 220
Db
       661 CGTTACAGTTGCACTCCCAGGAATTACTCGGTCAATATAAGAGAAGAGCTGAAGTTGGCC 720
Qу
          221 ArgTyrSerCysThrProArgAsnTyrSerValAsnIleArgGluGluLeuLysLeuAla 240
Db
       721 AATGTGGTCTTCTTCCACGTTGCCTCCTCGTGCAGCGCTGTGGAGGAAATTGTGGCTGT 780
Qу
          Db
       241 AsnValValPhePheProArgCysLeuLeuValGlnArgCysGlyGlyAsnCysGlyCys 260
       781 GGAACTGTCAACTGGAGGTCCTGCACATGCAATTCAGGGAAAACCGTGAAAAAGTATCAT 840
Qу
          Db
       261 GlyThrValAsnTrpArgSerCysThrCysAsnSerGlyLysThrValLysLysTyrHis 280
       841 GAGGTATTACAGTTTGAGCCTGGCCACATCAAGAGGGGGGGTAGAGCTAAGACCATGGCT 900
Qy
          Db
       281 GluValLeuGlnPheGluProGlyHisIleLysArgArgGlyArgAlaLysThrMetAla 300
       901 CTAGTTGACATCCAGTTGGATCACCATGAACGATGCGATTGTATCTGCAGCTCAAGACCA 960
Qу
         {\tt 301\ LeuValAspIleGlnLeuAspHisHisGluArgCysAspCysIleCysSerSerArgPro\ 320}
Db
       961 CCTCGA 966
Qy
          111111
Db
       321 ProArg 322
```

#### RESULT 2

US-09-457-066-37

<sup>;</sup> Sequence 37, Application US/09457066

<sup>;</sup> Patent No. 6432673